

Figure 2: Mouse Genome; One Mistmatch; Truncation to Length "L"

Theory and Measurements

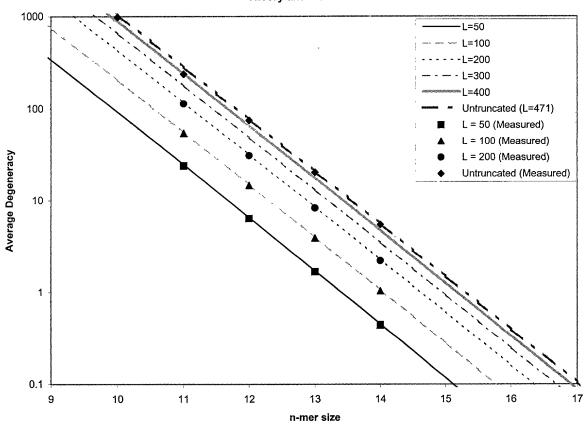
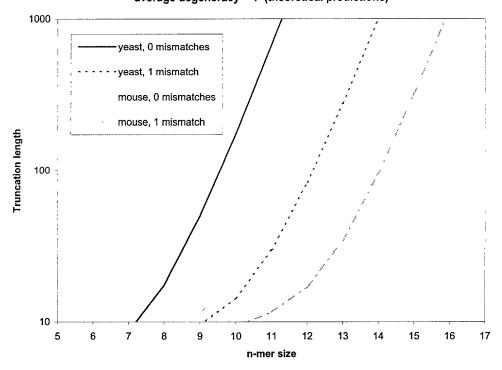
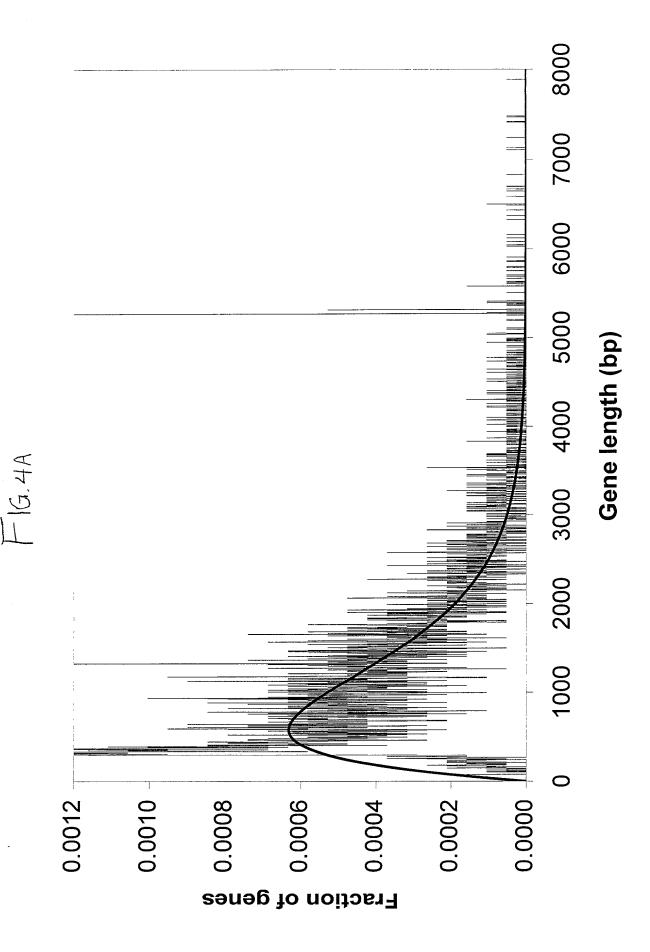
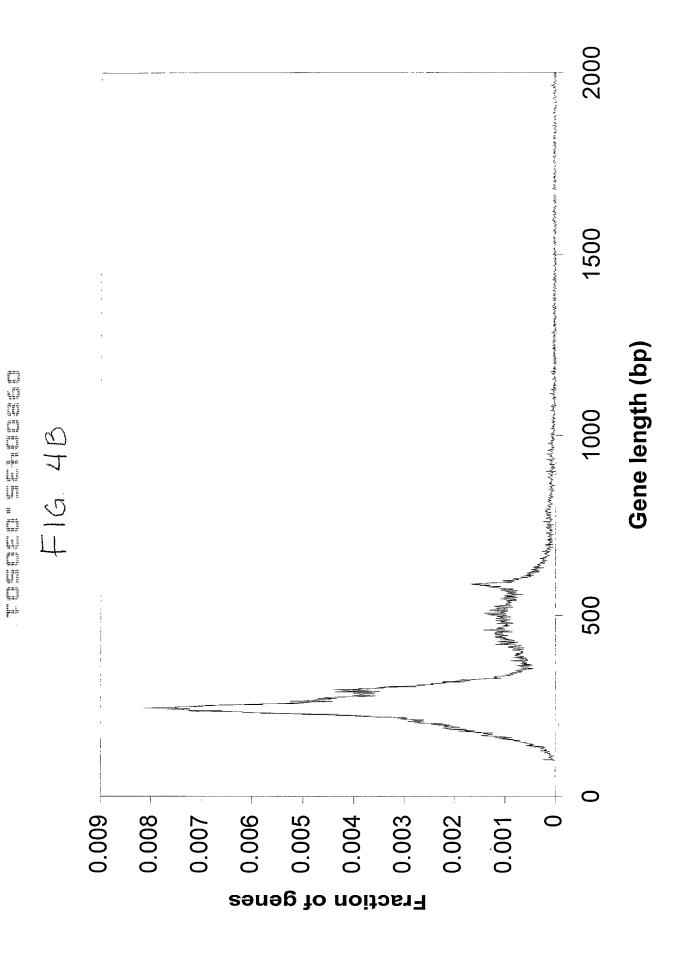
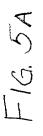


Figure 3: Relationships between parameter values such that average degeneracy = 1 (theoretical predictions)

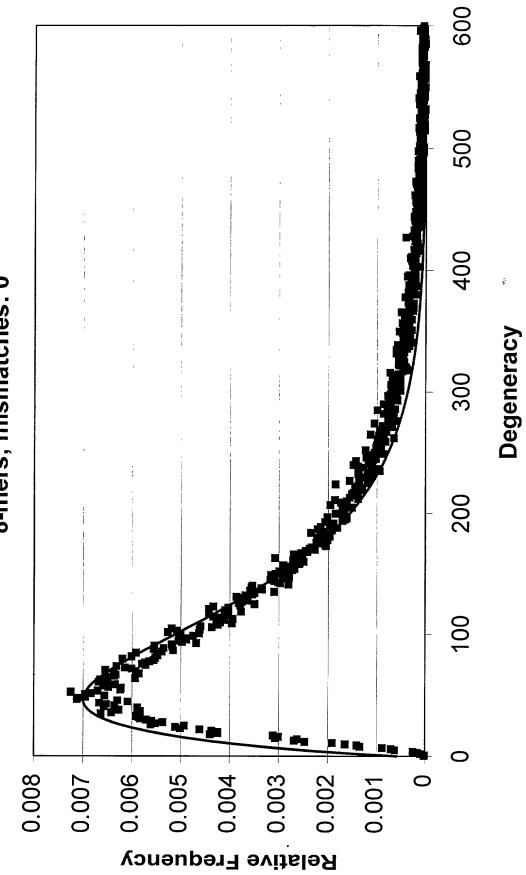


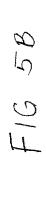




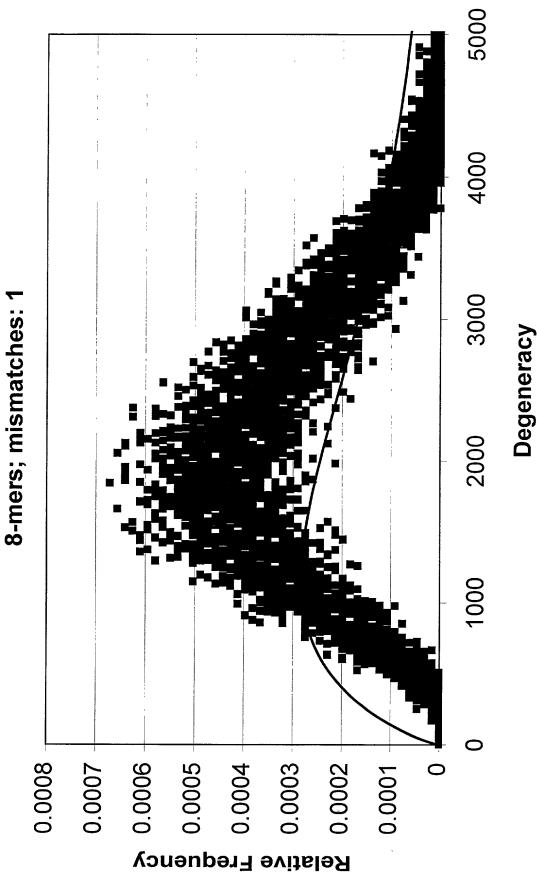


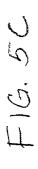




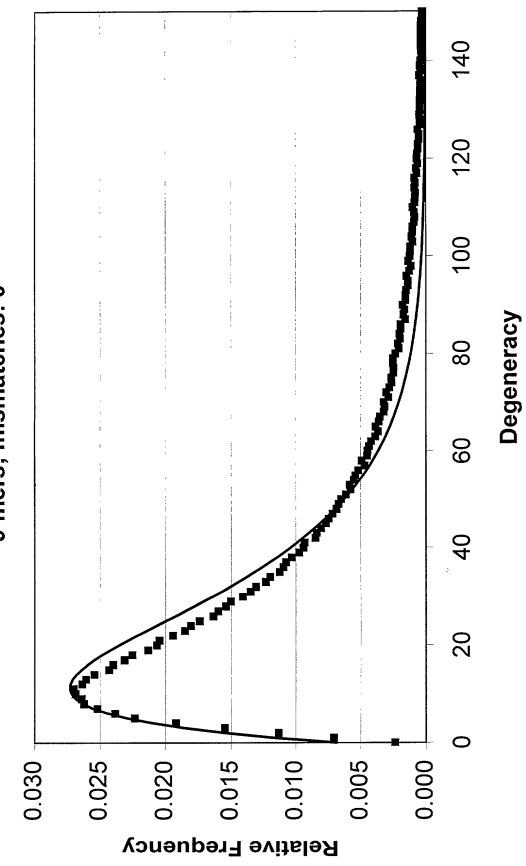


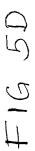




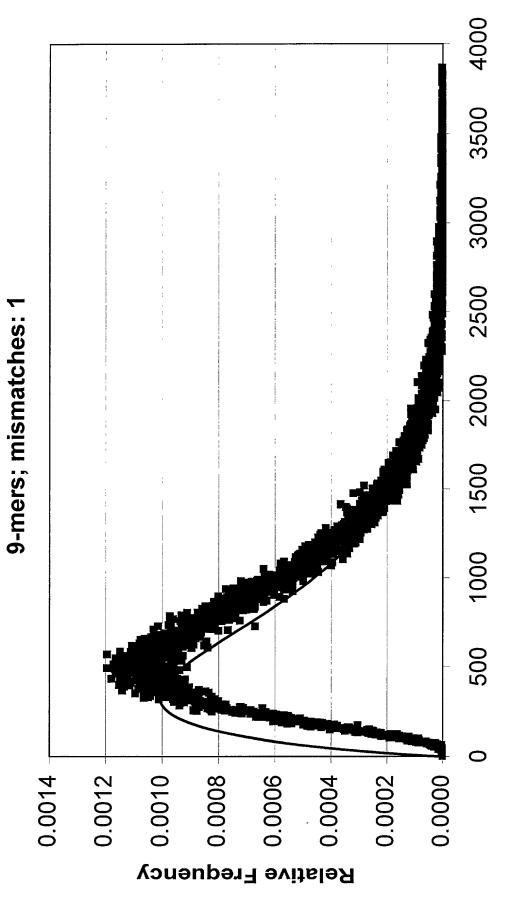


9-mers; mismatches: 0

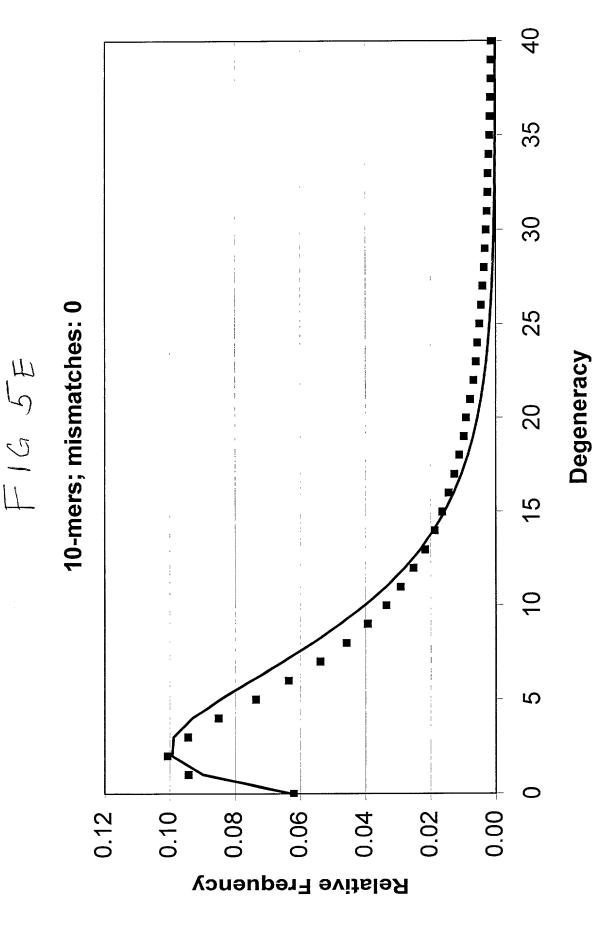




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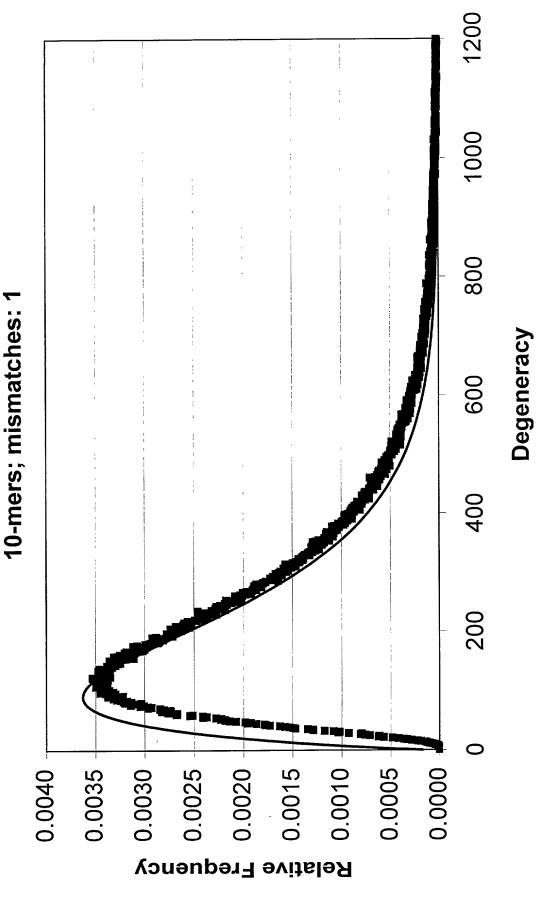


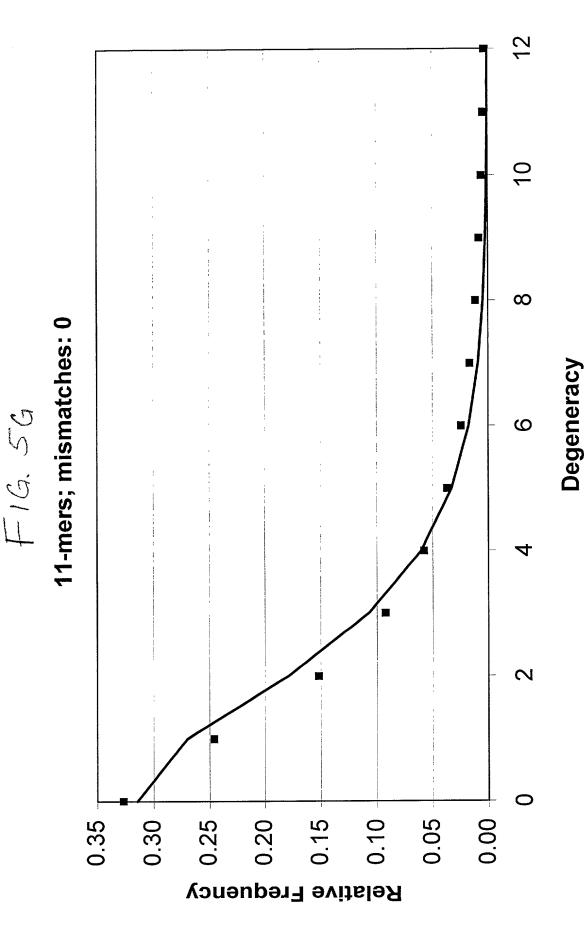
Degeneracy

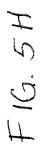




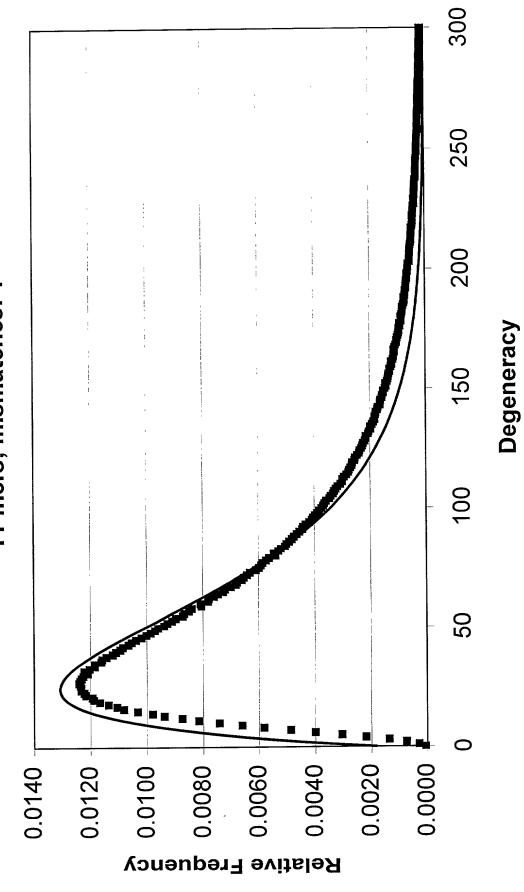




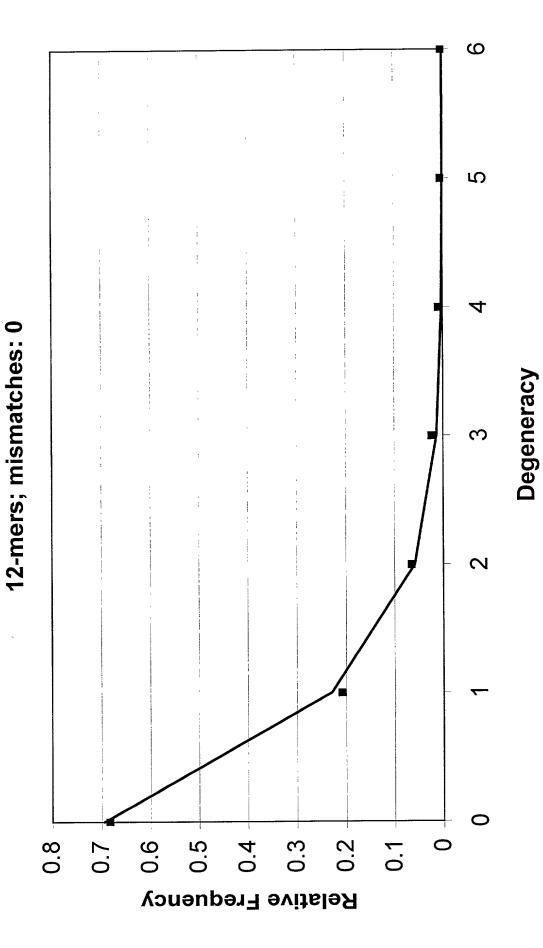




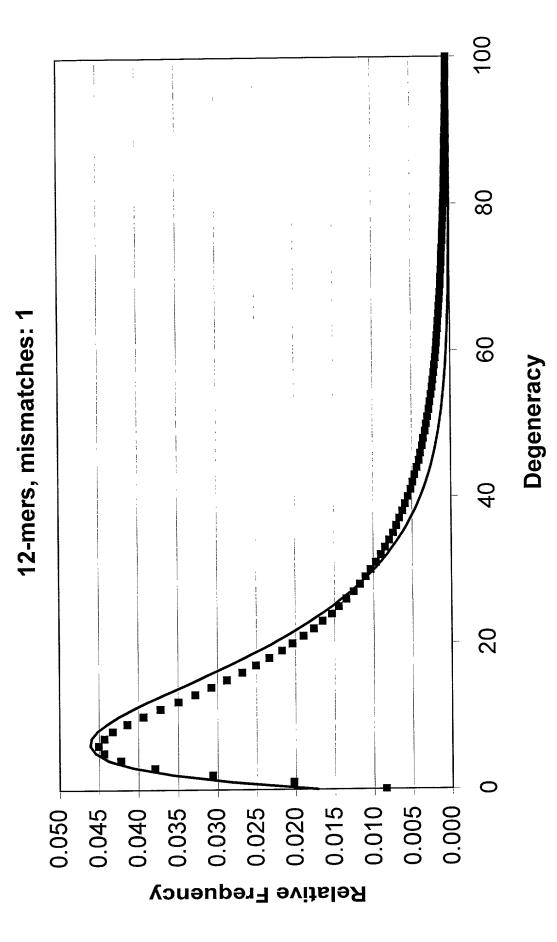
11-mers; mismatches: 1

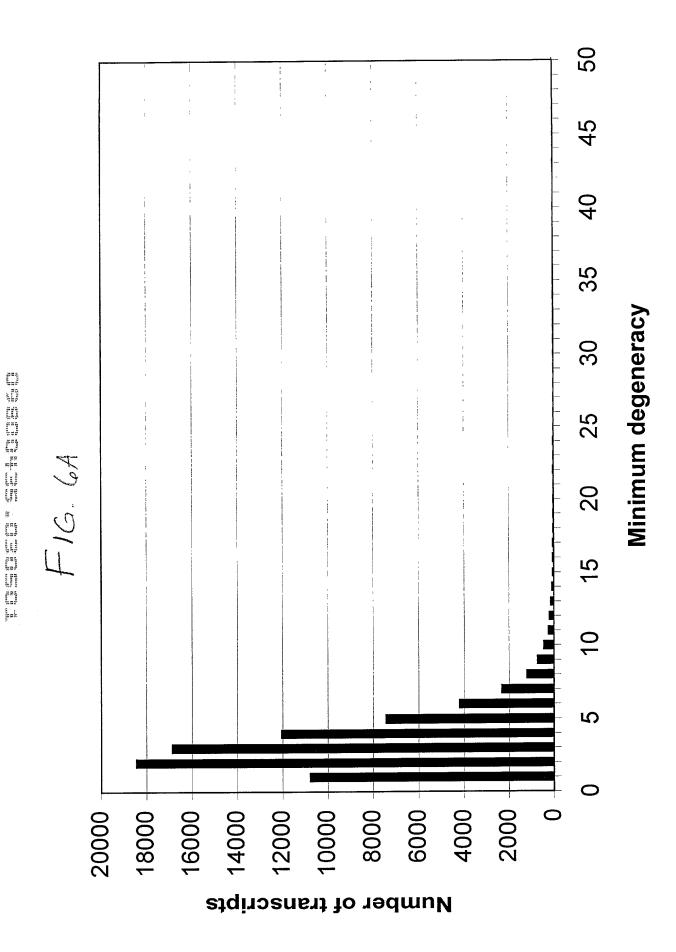


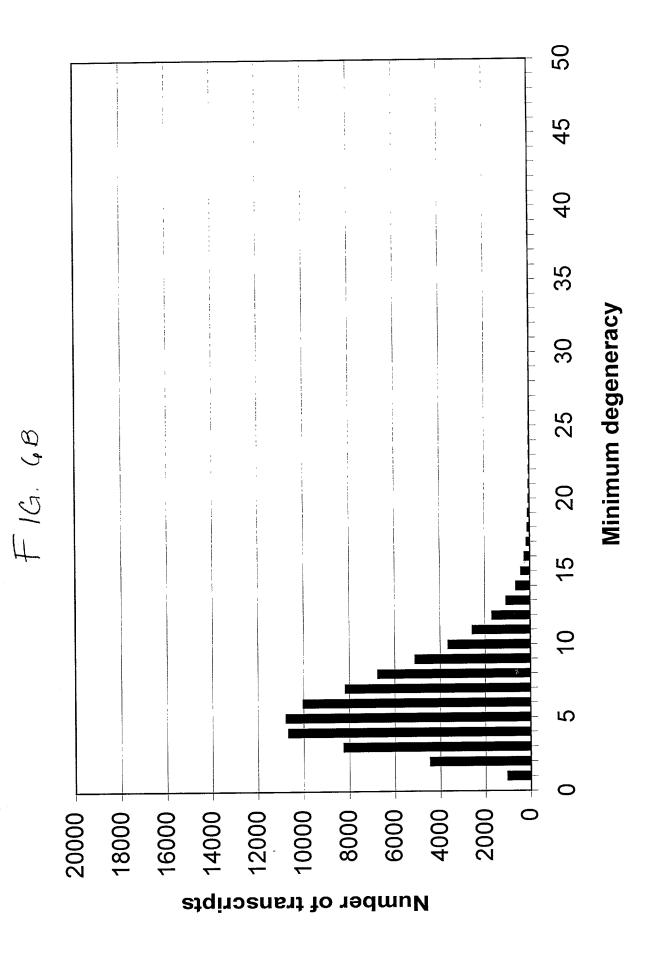
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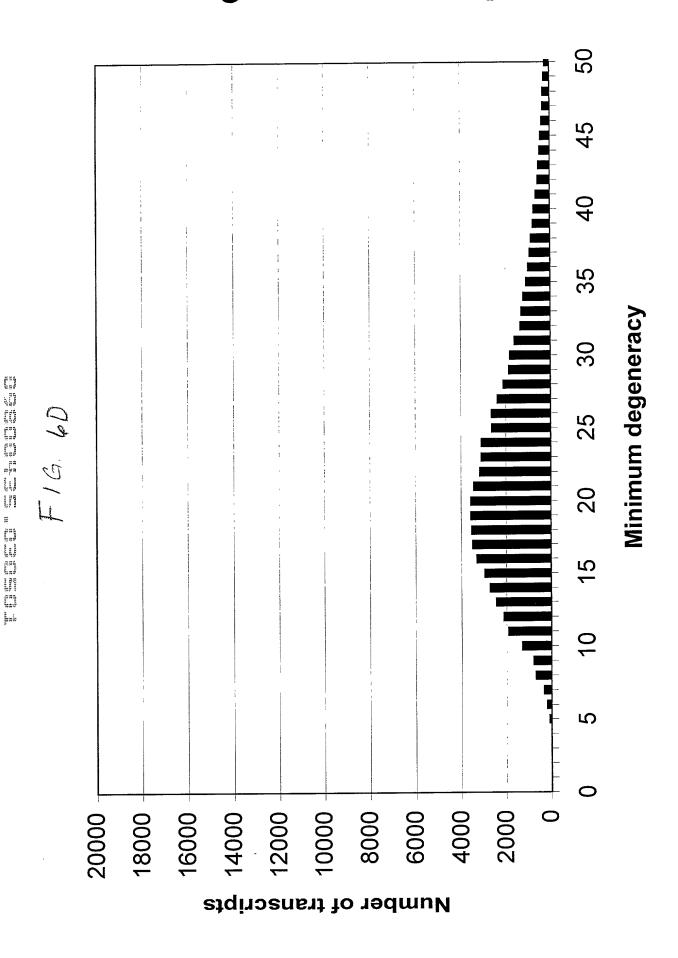


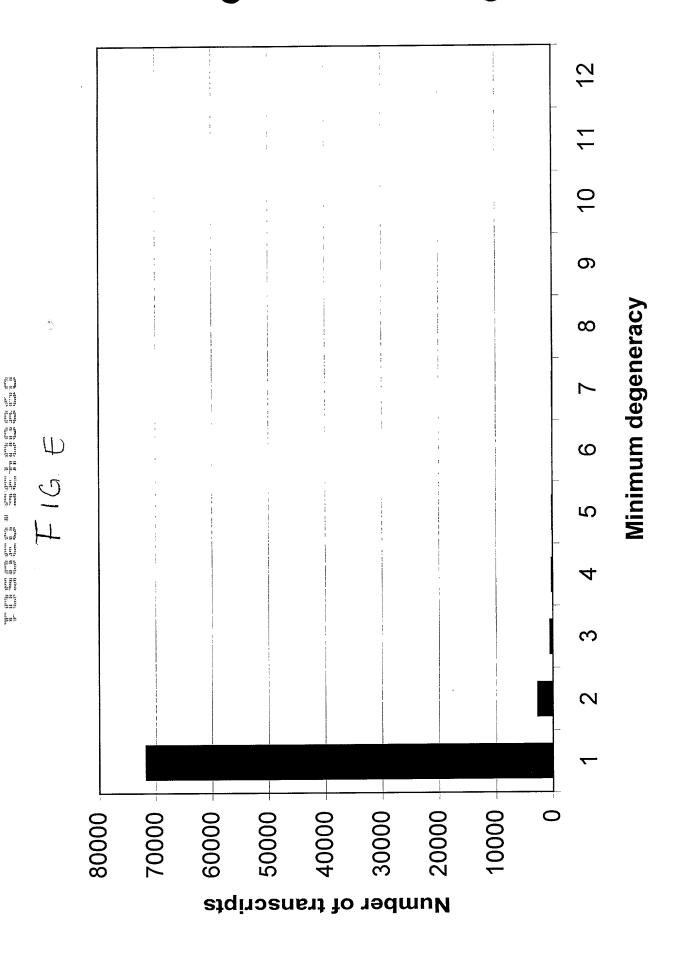


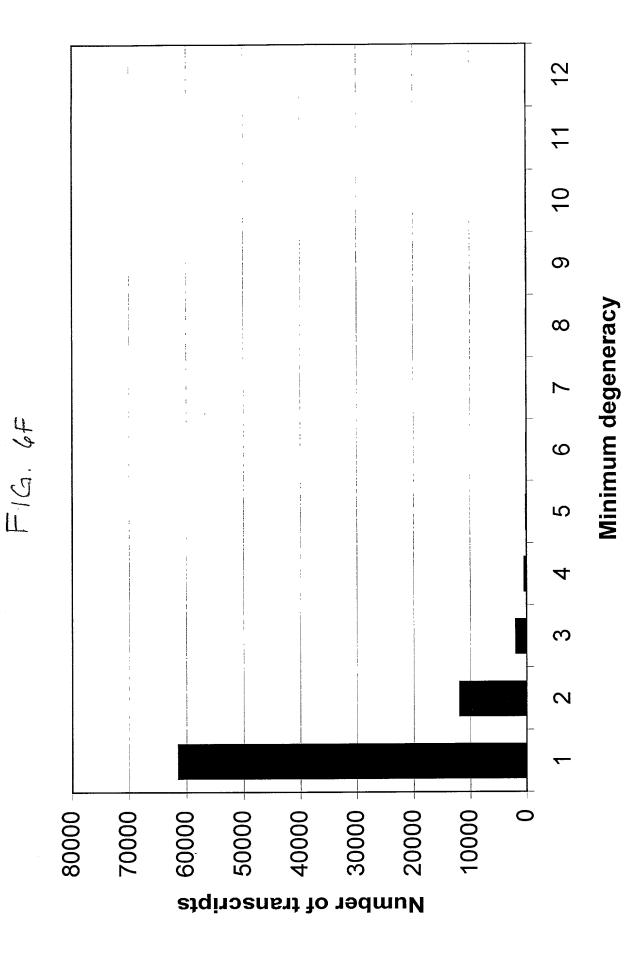




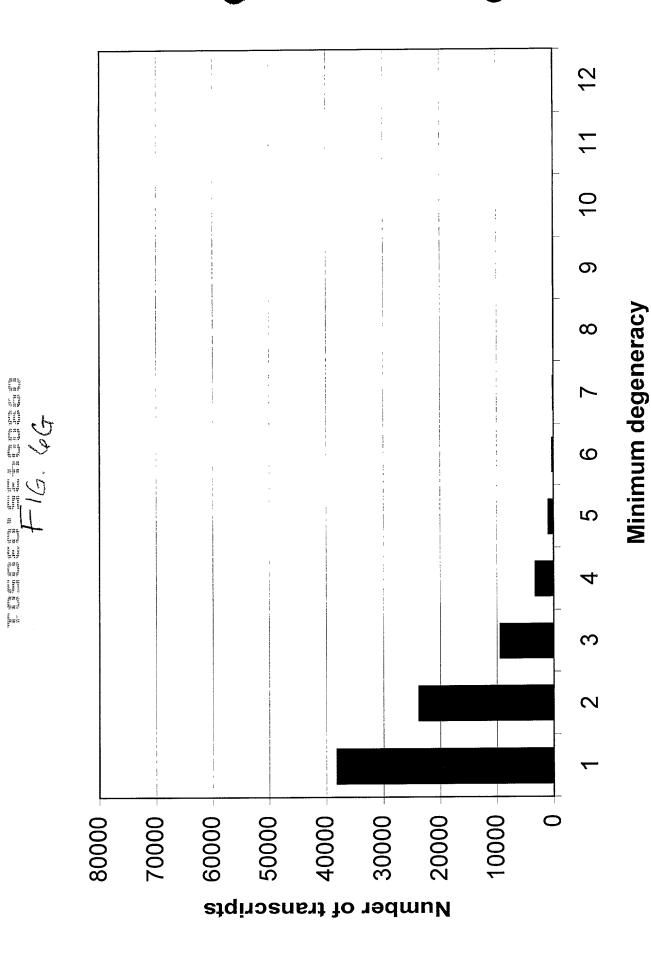
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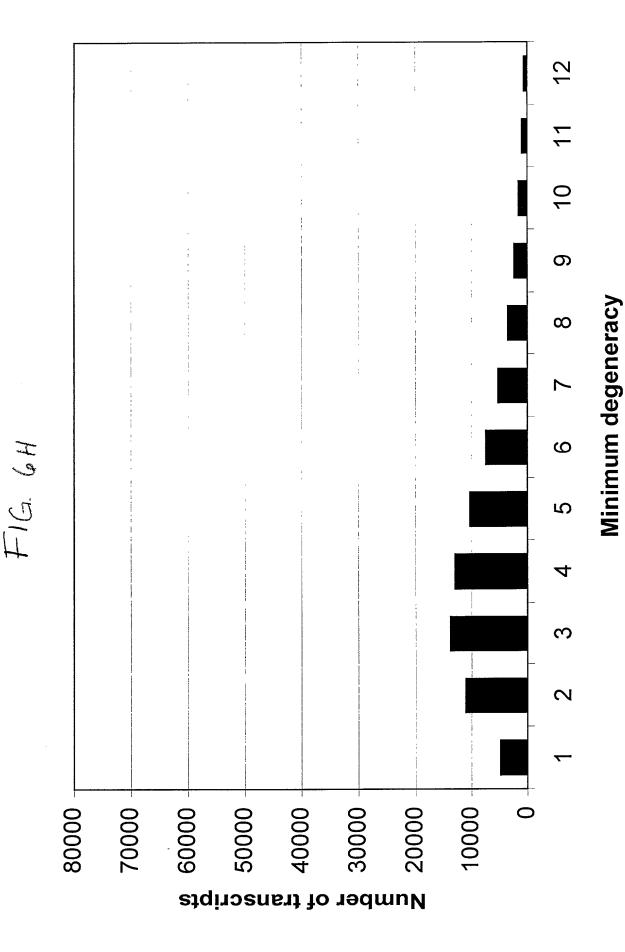




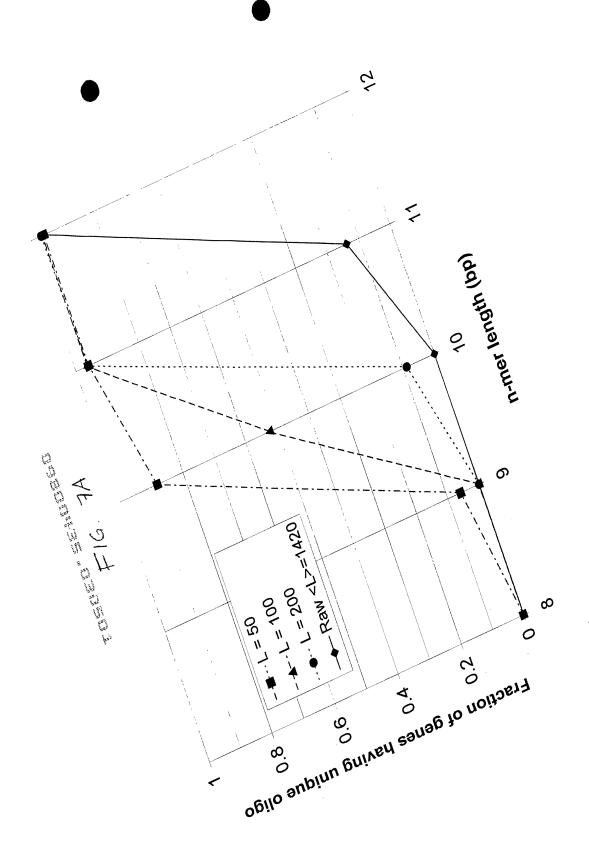


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